

Amendments to the Specification:

Please replace the paragraph beginning at page 28, line13 with the following amended paragraph:

Five regions of human genomic DNA that contain known single-nucleotide polymorphisms (SNPs) were PCR-amplified using tailed reverse primers to enable the formation of four-way complexes. The sequence of these regions, the location and identity of the respective SNPs within them and the sequences of the primers can be found in the National Center for Biotechnology Information (NCBI) SNP database (NCBI). ~~A database of single nucleotide polymorphisms shown at the web site of www.ncbi.nlm.nih.gov/SNP.~~ The NCBI accession IDs of the SNPs used were as follows: ss4215, ss4217, ss4213, ss4141, ss4212 and ss4030.